

SEQUENCE LISTING

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Neal Bringe
Katayoon Dehesh

<120> Nucleic Acid Constructs and Methods for Producing Altered Seed Oil Compositions

<130> 16518.133

<150> US 10/393,347
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<151> 2002-03-21

<150> US 60/390,185
<151> 2002-06-21

<160> 60

<170> PatentIn version 3.1

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<223> FAD3-1A intron 1

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<211> 184
<212> DNA
<213> Glycine max

<220>
<223> FAD3-1A 3'UTR

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agacatggaa tatttatttg aaattagtaa ggtagtaata ataaattttg aattgtcagt 180
ttca 184

<210> 17
<211> 143
<212> DNA
<213> Glycine max

<220>
<223> FAD3-1A 5'UTR

<400> 17
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tctaggtttt tacgcaccac gtatccctga gaaaagagag gaaccacact ctctaagcca 120
aagcaaaagc agcagcagca gca 143

<210> 18

<211> 2683
<212> DNA
<213> Glycine max

<220>
<223> partial FAD3-1B genomic clone

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<210> 19
<211> 160
<212> DNA
<213> Glycine max

<220>
<223> FAD3-1B intron 1

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tcagtgatgc tttagtcatt tcatttccact tggttatgca tgattggtcg ttcattatggt 120
ctgtcatggt gagttctaatt ttgattgatg catggaacag 160

<210> 20
<211> 119
<212> DNA
<213> Glycine max

<220>
<223> FAD3-1B intron 2

<400> 20
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<210> 21
<211> 166
<212> DNA
<213> Glycine max

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<223> FAD3-1B intron 3a

<400> 21
gtattactat gagtttgctt gattaatttc cacatttttt ctttcttctt aattttaatc 60
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ttcatgaagt ggttcatgat tatgtgtctt tatgccttta tgtcag 166

<210> 22
<211> 156
<212> DNA
<213> Glycine max

<220>
<223> FAD3-1B intron 3b

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tcactctctc tttgtgatat gaaccatata tttcag 156

<210> 23
<211> 148
<212> DNA
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<220>
<223> FAD3-1B intron 3c

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tcacttattt gaacttgaac atgtgtag 148

<210> 24
<211> 351
<212> DNA
<213> Glycine max

<220>
<223> FAD3-1B intron 4

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<210> 25
<211> 277
<212> DNA
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<220>
<223> FAD3-1B intron 5

<400> 25
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aataatttct cctattctac aatcaataat ccttctatgg tcctgaattg ttcctttctt 240
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<210> 26
<211> 158
<212> DNA
<213> Glycine max

<220>
<223> FAD3-1B 3'UTR

<400> 26

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 tgctagaat gcacttactt ttcaaagcat gctatgtc 158

<210> 27
 <211> 83
 <212> DNA
 <213> Glycine max

<220>
 <223> FAD3-1B 5'UTR

<400> 27
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<210> 28
 <211> 4083
 <212> DNA
 <213> Glycine max

<220>
 <223> FATB-1 genomic clone

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<212> DNA
<213> Glycine max

<220>
<223> FATB-1 intron I

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<210> 30
<211> 836
<212> DNA
<213> Glycine max

<220>
<223> FATB-1 intron II

<400> 30
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<212> DNA
<213> Glycine max

<220>
<223> FATB-1 intron III

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gttgttactt ttcatactat atttatatca actatttgc taacaacag 169

<210> 32
<211> 525
<212> DNA
<213> Glycine max

<220>
<223> FATB-1 intron IV

<400> 32
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<210> 33
<211> 389
<212> DNA
<213> Glycine max

<220>
<223> FATB-1 intron V

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389

<210> 34
<211> 106
<212> DNA
<213> Glycine max

<220>
<223> FATB-1 intron VI

<400> 34
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tggtgctttc taattagttt acattatgta tcttcattct tccagt 106

<210> 35
<211> 82
<212> DNA
<213> Glycine max

<220>
<223> FATB-1 intron VII

<400> 35
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actgtccttt gattgtttgc ag 82

<210> 36
<211> 208
<212> DNA
<213> Glycine max

<220>
<223> FATB-1 3'UTR

<400> 36
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<210> 37
<211> 229
<212> DNA
<213> Glycine max

<220>
<223> FATB-1 5'UTR

<400> 37
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<210> 38
<211> 1398
<212> DNA
<213> *Cuphea pulcherrima*

<220>
<223> KAS I gene

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<210> 39
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 <212> DNA
 <213> *Cuphea pulcherrima*

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<212> DNA
<213> Ricinus communis

<220>
<223> delta-9 desaturase

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<210> 41
<211> 1194
<212> DNA

<213> *Simmondsia chinensis*

<220>

<223> delta-9 desaturase

<400> 41

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<210> 42

<211> 2077

<212> DNA

<213> Artificial

<220>

<223> FATB-2 cDNA Contig

<400> 42

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<210> 43
 <211> 4634
 <212> DNA
 <213> Glycine max

<400> 43
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4634

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<211> 1215
<212> DNA
<213> Glycine max

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tccaactcct catataattc aagacaaaat cccgcgtttt ctgcatttct agacgttcta 180
ccctacaagg ttctcgattc ttcttttttc ttttttttta gactattatt attttaaaaa 240
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<210> 45
<211> 338
<212> DNA
<213> Glycine max

<400> 45
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 taattgcata ggacaaaact tacctacagt tcgtttgaca ttttttgtgt cgtttttaaa 180
 tcaaaattaa aattttatct tggtaatttg cagattatta gatacaactc caatttcgat 240
 caaagaacaa tgccaaaaac acctatggaa tctaagtttt gtgcaattgc ttattgatga 300
 ttttatttta ttgcctaaat tgtctgtttt ccaaacag 338

<210> 46
 <211> 641
 <212> DNA
 <213> Glycine max

<400> 46
 gtaagtgggt gtgactaaga agaacctttt tgatgtgtga agaattgcaa aggcgtccat 60
 gtcagctgt gaaatcttct tttgccttac tcatctttac tttgacttta tatagtatct 120
 ggttgaatta ttttgactt ctgcatttgt ttctgtcact tgtgcttttt tgtttcacia 180
 aattggatg atagtttaga acttgggatt aaaggcatgt ttggaatata ttgtgattgt 240
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 gaataaaaaa taaaactact gtaatgtgta taaaaaattc ttcttgatg gtaattgatc 360
 tgataagcac atgcttttta cataatgaat tatatgaagt cttttgcctt aagtctgtta 420
 gactgggtat gagatatggt agtaaattct ttttacatc cgtacatttt tttgcatatt 480
 tctgtcttat tattgtaaaa tgttggatgc atatacaggt tttcaaaaga agcaacttat 540
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<210> 47
 <211> 367
 <212> DNA
 <213> Glycine max

<400> 47
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 tgtgaagtta tctattatca tatttacttt ctcttaatac accactattg aaaggcaatt 180
 cattacagat ttaagcatac aaaattttgt tgatgataat tttttaatct accaacagta 240

tctaatatct tcttaatttg ttattaagta ccagccttca acttgtgtac atgttgacc 300
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<211> 18
<212> DNA
<213> Artificial sequence

<220>
<223> PCR primer

<400> 48 18
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<210> 49
<211> 17
<212> DNA
<213> Artificial sequence

<220>
<223> PCR primer

<400> 49 17
cttctcgttt gttgagc

<210> 50
<211> 16
<212> DNA
<213> Artificial sequence

<220>
<223> PCR primer

<400> 50 16
cagctgcaac ttcac

<210> 51
<211> 16
<212> DNA
<213> Artificial sequence

<220>
<223> PCR primer

<400> 51 16
cttccccatt aggtcc

<210> 52

<211> 18
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<220>
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<400> 52
cacttaatca tgттаага 18

<210> 53
<211> 17
<212> DNA
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<220>
<223> PCR primer

<400> 53
gtcgtgattg gcttgтg 17

<210> 54
<211> 17
<212> DNA
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<220>
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<400> 54
ctctgctcca gttgtgc 17

<210> 55
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<220>
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<400> 55
gcgagggtga agтаacag 18

<210> 56
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<212> DNA
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<220>
<223> PCR primer

<400> 56
gcacaaacct tgттtctg 18

<210> 57
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<220>
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<400> 57
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17

<210> 58
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<212> DNA
<213> Artificial sequence

<220>
<223> PCR primer

<400> 58
gatttcacca gatttcg

17

<210> 59
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<212> DNA
<213> Artificial sequence

<220>
<223> PCR primer

<400> 59
gtgcgaatga aattagc

17

<210> 60
<211> 17
<212> DNA
<213> Artificial sequence

<220>
<223> PCR primer

<400> 60
ctttctgctg gaactgg

17